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AUG 0 9 2001

TECH CENTER 1600/2900

SEQUENCE LISTING

<110> Seguela, Philippe Babinski, Kazimierz McGill University

<120> DNA ENCODING A HUMAN PROTON-GATED ION CHANNEL AND USES THEREOF

<130> 641050.90021

<140> CA PCT/CA98/01016

<141> 1998-10-29

<150> US09/530,233

<151> 1997-10-29

<160> 5

<170> PatentIn Ver. 2.0

<210> 1

<211> 1732

<212> DNA

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Arg Arg Gln Pro Ser Asp Ile Arg Val Phe Ala Ser Asn Cys Ser Met

cac ggg ctg ggc cac gtc ttc ggg cca ggc agc ctg agc ctg cgc cgg 147 His Gly Leu Gly His Val Phe Gly Pro Gly Ser Leu Ser Leu Arg Arg 30 35 40

ggg atg tgg gca gcg gcc gtg gtc ctg tca gtg gcc acc ttc ctc tac 195 Gly Met Trp Ala Ala Ala Val Val Leu Ser Val Ala Thr Phe Leu Tyr 45 50 55

cag gtg gct gag agg gtg cgc tac tac agg gag ttc cac cac cag act 243

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•		GIN	Va1 60	нта	GIU	Arg	val	Arg 65	ıyr	ıyr	Arg	GIU	Phe 70	nlS	нlS	GIN	ınr	
													. •					
		_	_	_		cga	_	_				_		_	-	-		291
	,		Leu	Asp	Glu	Arg		Ser	His	Arg	Leu		Phe	Pro	Ala	Val		
		75					80					85					90	
		ctg	tgc	aac	atc	aac	cca	ctg	cgc	cgc	tcg	cgc	cta	acg	ccc	aac	gac	339
		_	-			Asn		_	_	_	_	_		_			_	
						95					100					105		
		cta	Cac	taa	act	ggg	+ < +	aca	cta	cta	aac	cta	αa+	ccc	ac =	nan	Cac	387
		_			_	Gly		-					-					507
			-	_	110	-				115	_		-		120			
		_	_			cgc	_	_					_					435
		Ala	АТа	125	ьeu	Arg	Ala	ьeu	130	Arg	PTO	FIO	ATa	135	FTO	σтλ	rne	
	-																	
		_		_		acc												483
	•	Met		Ser	Pro	Thr	Phe	_	Met	Ala	Gln	Leu	_	Ala	Arg	Ala	Gly	
			140					145					150					
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		tat	gaa	cct	gan	aac	tta	acc	aca	atc	ttc	acc	caa	ato	gga	aaa	tac	579
		•				Asn			-									
	•					175					180					185		
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		Glu		Tyr	Leu	Pro	Val		Arg	Asp	Asn	Glu		Thr	Pro	Phe	Glu	
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		235					240					245					250	
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_	cgc Arg	_	_	_	_		-			_		_	_		-	1011
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tac . Tyr	-	-		-												14
cga Arg 475	_															14
ctg Leu																15
cct Pro																15
cgc Arg			tac				Gln		taga	acct	gct (gtctq	gtgto	cc		16
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tegg cccc		ccc ç					acat				acgt	cagct	itt t	cegt	cttca	
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Ser His Arg Leu Val Phe Pro Ala Val Thr Leu Cys Asn Ile Asn Pro Leu Arg Arg Ser Arg Leu Thr Pro Asn Asp Leu His Trp Ala Gly Ser Ala Leu Leu Gly Leu Asp Pro Ala Glu His Ala Ala Phe Leu Arg Ala Leu Gly Arg Pro Pro Ala Pro Pro Gly Phe Met Pro Ser Pro Thr Phe Asp Met Ala Gln Leu Tyr Ala Arg Ala Gly His Ser Leu Asp Asp Met Leu Leu Asp Cys Arg Phe Arg Gly Gln Pro Cys Gly Pro Glu Asn Phe Thr Thr Ile Phe Thr Arg Met Gly Lys Cys Tyr Thr Phe Asn Ser Gly Ala Asp Gly Ala Glu Leu Leu Thr Thr Arg Gly Gly Met Gly Asn Gly Leu Asp Ile Met Leu Asp Val Gln Glu Glu Tyr Leu Pro Val Trp Arg Asp Asn Glu Glu Thr Pro Phe Glu Val Gly Ile Arg Val Gln Ile His Ser Gln Glu Glu Pro Pro Ile Ile Asp Gln Leu Gly Leu Gly Val Ser Pro Gly Tyr Gln Thr Phe Val Ser Cys Gln Gln Gln Leu Ser Phe Leu Pro Pro Pro Trp Gly Asp Cys Ser Ser Ala Ser Leu Asn Pro Asn Tyr Glu Pro Glu Pro Ser Asp Pro Leu Gly Ser Pro Ser Pro Ser Pro Ser Pro Pro Tyr Thr Leu Met Gly Cys Arg Leu Ala Cys Glu Thr Arg Tyr Val Ala Arg Lys Cys Gly Cys Arg Met Val Tyr Met Pro

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Pro Ser Arg Ala Ala Ala Arg Phe Leu Ala Arg Lys Leu Asn Arg Ser 385 390 395 400

Glu Ala Tyr Ile Ala Glu Asn Val Leu Ala Leu Asp Ile Phe Phe Glu 405 410 415

Ala Leu Asn Tyr Glu Thr Val Glu Gln Lys Lys Ala Tyr Glu Met Ser 420 425 430

Glu Leu Leu Gly Asp Ile Gly Gly Gln Met Gly Leu Phe Ile Gly Ala 435 440 445

Ser Leu Leu Thr Ile Leu Glu Ile Leu Asp Tyr Leu Cys Glu Val Phe 450 455 460

Arg Asp Lys Val Leu Gly Tyr Phe Trp Asn Arg Gln His Ser Gln Arg 465 470 475 480

His Ser Ser Thr Asn Leu Leu Gln Glu Gly Leu Gly Ser His Arg Thr 485 490 495

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